

## STIC Biotechnology Systems Branch

### RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 10/543,033  
Source: Put/10  
Date Processed by STIC: 7/28/05

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 4.2.2 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

<http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm>

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.

Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<http://www.uspto.gov/ebc/efs/downloads/documents.htm>), EFS Submission User Manual - ePAVE)
2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
3. Hand Carry, Federal Express, United Parcel Service, or other delivery service (EFFECTIVE 01/14/05): U.S. Patent and Trademark Office, Mail Stop Sequence, Customer Window, Randolph Building, 401 Dulany Street, Alexandria, VA 22314

Revised 01/24/05

## Raw Sequence Listing Error Summary

**ERROR DETECTED**
**SUGGESTED CORRECTION**
**SERIAL NUMBER:** 10/543,033

**ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE**

1  Wrapped Nucleic  
Wrapped Aminos      The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor after creating it. Please adjust your right margin to .3; this will prevent "wrapping."

2  Invalid Line Length      The rules require that a line **not exceed** 72 characters in length. This includes white spaces.

3  Misaligned Amino  
Numbering      The numbering under each 5<sup>th</sup> amino acid is misaligned. **Do not** use tab codes between numbers; use **space characters**, instead.

4  Non-ASCII      The submitted file was **not saved** in ASCII(DOS) text, as **required** by the Sequence Rules. **Please ensure your subsequent submission is saved in ASCII text.**

5  Variable Length      Sequence(s) \_\_\_\_\_ contain n's or Xaa's representing more than one residue. **Per Sequence Rules, each n or Xaa can only represent a single residue.** Please present the **maximum** number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.

6  PatentIn 2.0  
"bug"      A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequences(s) \_\_\_\_\_. Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. **This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences.**

7  Skipped Sequences  
(OLD RULES)      Sequence(s) \_\_\_\_\_ missing. If intentional, please insert the following lines for **each** skipped sequence:  
(2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)  
(i) SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading)  
(xi) SEQUENCE DESCRIPTION:SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)  
This sequence is intentionally skipped  
  
Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to **include** the skipped sequences.

8  Skipped Sequences  
(NEW RULES)      Sequence(s) \_\_\_\_\_ missing. If **intentional**, please insert the following lines for **each** skipped sequence.  
<210> sequence id number  
<400> sequence id number  
000

9  Use of n's or Xaa's  
(NEW RULES)      Use of n's and/or Xaa's have been detected in the Sequence Listing.  
Per 1.823 of Sequence Rules, use of <220>-<223> is MANDATORY if n's or Xaa's are present.  
In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.

10  Invalid <213>  
Response      Per 1.823 of Sequence Rules, the only **valid** <213> responses are: Unknown, Artificial Sequence, or scientific name (Genus/species). <220>-<223> section is **required** when <213> response is Unknown or is Artificial Sequence

11  Use of <220>  
  
→      Sequence(s) \_\_\_\_\_ missing the <220> "Feature" and associated numeric identifiers and responses.  
Use of <220> to <223> is MANDATORY if <213> "Organism" response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in <220> to <223> section.  
(See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of Sequence Rules)

12  PatentIn 2.0  
"bug"      Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.

13  Misuse of n/Xaa      "n" can **only** represent a single nucleotide; "Xaa" can **only** represent a single amino acid



PCT

## RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/543,033

DATE: 07/28/2005

TIME: 17:04:05

Input Set : D:\seq listing 10589-012-999 (as filed).txt  
 Output Set: N:\CRF4\07282005\J543033.raw

3 <110> APPLICANT: Cao, Liangxian  
 4 Trifillis, Panayiota  
 6 <120> TITLE OF INVENTION: METHODS FOR IDENTIFYING COMPOUNDS THAT MODULATE UNTRANSLATED  
 7 REGION-DEPENDENT GENE EXPRESSION AND METHODS OF USING SAME  
 9 <130> FILE REFERENCE: 10589-012-999  
 C--> 11 <140> CURRENT APPLICATION NUMBER: US/10/543,033  
 C--> 12 <141> CURRENT FILING DATE: 2005-07-21  
 14 <150> PRIOR APPLICATION NUMBER: PCT/US2004/001643  
 15 <151> PRIOR FILING DATE: 2004-01-21  
 17 <150> PRIOR APPLICATION NUMBER: 60/441,637  
 18 <151> PRIOR FILING DATE: 2003-01-21  
 20 <160> NUMBER OF SEQ ID NOS: 94  
 22 <170> SOFTWARE: PatentIn version 3.2  
 24 <210> SEQ ID NO: 1  
 25 <211> LENGTH: 14  
 26 <212> TYPE: DNA  
 27 <213> ORGANISM: Artificial Sequence  
 29 <220> FEATURE:  
 30 <223> OTHER INFORMATION: Description of Artificial Sequence: one motif of G-quartet  
 element  
 33 <220> FEATURE:  
 34 <221> NAME/KEY: misc\_feature  
 35 <222> LOCATION: 3, 7, 8, 11  
 36 <223> OTHER INFORMATION: n = a, t, c, or g  
 38 <220> FEATURE:  
 39 <221> NAME/KEY: misc\_feature  
 40 <222> LOCATION: (7)..(8)  
 41 <223> OTHER INFORMATION: This represents one form of the sequence as described, other forms  
 42 described may have up to five nucleotides in this variable region  
 44 <400> SEQUENCE: 1  
 W--> 45 ggntgggnngg ntgg  
 48 <210> SEQ ID NO: 2  
 49 <211> LENGTH: 14  
 50 <212> TYPE: DNA  
 51 <213> ORGANISM: Artificial Sequence  
 53 <220> FEATURE:  
 54 <223> OTHER INFORMATION: Description of Artificial Sequence: one motif of G-quartet  
 element  
 57 <220> FEATURE:  
 58 <221> NAME/KEY: misc\_feature  
 59 <222> LOCATION: 3, 4, 7, 8, 11, 12  
 60 <223> OTHER INFORMATION: n = a, t, g or c  
 62 <220> FEATURE:

*ppr 1,3,5-8*  
*Does Not Comply*  
*Corrected Diskette Nocde*

*give source of genetic material*

*one motif of G-quartet*

*( see item 11 on Error*  
*summary sheet)*

14

63 <221> NAME/KEY: misc\_feature  
64 <222> LOCATION: (2)..(12)

## RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/543,033

DATE: 07/28/2005

TIME: 17:04:05

Input Set : D:\seq listing 10589-012-999 (as filed).txt  
 Output Set: N:\CRF4\07282005\J543033.raw

65 <223> OTHER INFORMATION: This represents one form of the sequence as described, other forms

66 described have longer variable regions, typical is 2 - 10  
 67 nucleotides

69 <400> SEQUENCE: 2

W--> 70 ggnnggnngg nngg

14

73 <210> SEQ ID NO: 3

74 <211> LENGTH: 14

75 <212> TYPE: DNA

76 <213> ORGANISM: Artificial Sequence

78 <220> FEATURE:

79 <223> OTHER INFORMATION: Description of Artificial Sequence: general formula of G-quartet élément

82 <220> FEATURE:

83 <221> NAME/KEY: misc\_feature

84 <222> LOCATION: 3, 4, 7, 8, 11, 12

85 <223> OTHER INFORMATION: n = a, t, g, or c

87 <220> FEATURE:

88 <221> NAME/KEY: misc\_feature

89 <222> LOCATION: (2)..(12)

90 <223> OTHER INFORMATION: This represents one form of the sequence as described, other forms

91 described have longer variable regions, typical is 2 - 10  
 92 nucleotides

94 <400> SEQUENCE: 3

14

W--> 95 ggnnggnngg nngg

98 <210> SEQ ID NO: 4

99 <211> LENGTH: 19

100 <212> TYPE: RNA

101 <213> ORGANISM: Artificial Sequence

103 <220> FEATURE:

104 <223> OTHER INFORMATION: Description of Artificial Sequence: one subunit of 15-LOX-DICE

106 <400> SEQUENCE: 4

19

107 ccccrccuc uucccaag

110 <210> SEQ ID NO: 5

111 <211> LENGTH: 152

112 <212> TYPE: DNA

113 <213> ORGANISM: Homo sapiens

115 <400> SEQUENCE: 5

60

116 gcagaggacc agctaagagg gagagaagca actacagacc cccccctgaaa acaaccctca

120

118 gacgccacat cccctgaccaa gctgccagc aggttctctt cctctcacat actgaccac

152

120 ggctccaccc tctctcccct ggaaaggaca cc

123 <210> SEQ ID NO: 6

124 <211> LENGTH: 792

125 <212> TYPE: DNA

126 <213> ORGANISM: Homo sapiens

128 <400> SEQUENCE: 6

60

129 ttaggaggac gaacatccaa ctttccccaa cgccctccct gccccaaatcc ctttattacc

120

131 ccctccctca gacaccctca acctcttctg gctaaaaag agaattgggg gcttagggtc

180

133 ggaacccaag cttagaaactt taagcaacaa gaccaccact tcgaaacctg ggattcagg

135 atgtgtggcc tgcacagtga attgctggca accactaaga attcaaactg gggctccag

240

137 aactcactgg ggcctacagc tttatccct gacatctgga atctggagac caggagcct

300

## RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/543,033

DATE: 07/28/2005

TIME: 17:04:05

Input Set : D:\seq listing 10589-012-999 (as filed).txt  
 Output Set: N:\CRF4\07282005\J543033.raw

139	ttggttctgg ccagaatgct gcaggacttg agaagacctc acctagaaat tgacacaagt	360
141	ggaccttagg ccttcctctc tccagatgtt tccagacttc cttgagacac ggagcccagc	420
143	cctccccatg gagccagctc cctctattta tgtttgcact tgtgattatt tattatttat	480
145	ttattattta ttatattaca gatgaatgta ttatattggg agaccgggt atcctgggg	540
147	acccaatgta ggagctgcct tggctcagac atgtttccg taaaaacgga gctgaacaat	600
149	agctgttcc catgtageccc cctggcctct gtgccttctt ttgattatgt tttttaaat	660
151	atttatctga ttaagttgtc taaaacaatgc tgatttggg accaactgtc actcattgct	720
153	gagcctctgc tccccagggg agttgtgtct gtaatcgccc tactattcag tggcgagaaa	780
155	taaagtttgc tt	792
158	<210> SEQ ID NO: 7	
159	<211> LENGTH: 21	
160	<212> TYPE: RNA	
161	<213> ORGANISM: Artificial Sequence	
163	<220> FEATURE:	
164	<223> OTHER INFORMATION: Description of Artificial Sequence: Group I AU-Rich element	
(ARE)		
165	cluster of 3'untranslated region	
167	<400> SEQUENCE: 7	
168	auuuauuuau uuaauuuauuu a	21
171	<210> SEQ ID NO: 8	
172	<211> LENGTH: 40	
173	<212> TYPE: DNA	
174	<213> ORGANISM: Homo sapiens	
176	<400> SEQUENCE: 8	
177	kctggaggat gtggctgcag agcctgctgc tcttgggcac	40
180	<210> SEQ ID NO: 9	
181	<211> LENGTH: 289	
182	<212> TYPE: DNA	
183	<213> ORGANISM: Homo sapiens	
185	<400> SEQUENCE: 9	
186	gccggggagc tgctctctca taaaacaaga gctagaaaact caggatggtc atcttggagg	60
188	gaccaagggg tggccacag ccatggtggg agtggcctgg acctgcctg gcccacactg	120
190	accctgatac aggcatggca gaagaatggg aatattttat actgacagaa atcagtaata	180
192	tttatatatatt tatatttta aaatattttat ttatttattt atttaagttc atattccata	240
194	tttattcaag atgtttacc gtaataatta ttataaaaa tatgcttct	289
197	<210> SEQ ID NO: 10	
198	<211> LENGTH: 21	
199	<212> TYPE: RNA	
200	<213> ORGANISM: Artificial Sequence	
202	<220> FEATURE:	
203	<223> OTHER INFORMATION: Description of Artificial Sequence: Group I AU-Rich element	
(ARE)		
204	cluster of 3'untranslated region	
206	<400> SEQUENCE: 10	
207	auuuauuuau uuaauuuauuu a	21
210	<210> SEQ ID NO: 11	
211	<211> LENGTH: 47	
212	<212> TYPE: DNA	
213	<213> ORGANISM: Homo sapiens	
215	<400> SEQUENCE: 11	
216	atcactctct ttaatcacta ctcacattaa cctcaactcc tgccaca	47

## RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/543,033

DATE: 07/28/2005

TIME: 17:04:05

Input Set : D:\seq listing 10589-012-999 (as filed).txt  
 Output Set: N:\CRF4\07282005\J543033.raw

219 <210> SEQ ID NO: 12  
 220 <211> LENGTH: 307  
 221 <212> TYPE: DNA  
 222 <213> ORGANISM: Homo sapiens  
 224 <400> SEQUENCE: 12

225 taattaatgtt	cttcccaactt	aaaacatatac	aggccttcta	tttatatttatt	taaatattta	60
227 aattttatat	ttatttgttga	atgtatgggtt	gctacctatt	gtaactatta	ttcttaatct	120
229 taaaactata	aatatggatc	ttttatgatt	ctttttgtaa	gcccttagggg	ctctaaaatg	180
231 gtttacctta	tttatccccaa	aaatatattt	tattatgtt	aatgttaat	atagtatcta	240
233 ttagatgtt	tttagaaaaac	tatthaataa	atttgataaa	tataaaaaaaa	aaaaacaaaaa	300
235 aaaaaaaaaa						307

238 <210> SEQ ID NO: 13  
 239 <211> LENGTH: 15  
 240 <212> TYPE RNA  
 241 <213> ORGANISM: Artificial Sequence  
 243 <220> FEATURE:  
 244 <223> OTHER INFORMATION: Description of Artificial Sequence: Group III AU-Rich element (ARE)

245 cluster of 3'untranslated region  
 248 <220> FEATURE:  
 249 <221> NAME/KEY: misc\_feature  
 250 <222> LOCATION: (1)..(15)  
 251 <223> OTHER INFORMATION: n = a, t, g or c      *It's not allowed in an RNA sequence.*  
 253 <400> SEQUENCE: 13

W--> 254 nauuuauuu uuuuu  
 257 <210> SEQ ID NO: 14  
 258 <211> LENGTH: 62  
 259 <212> TYPE: DNA  
 260 <213> ORGANISM: Homo sapiens  
 262 <400> SEQUENCE: 14

263 ttctgccttc	gagccacccg	ggaacgaaag	agaagctcta	tctcgccccc	aggagccca	60
265 ct						62

268 <210> SEQ ID NO: 15  
 269 <211> LENGTH: 427  
 270 <212> TYPE: DNA  
 271 <213> ORGANISM: Homo sapiens  
 273 <400> SEQUENCE: 15

274 tagcatgggc	acctcagatt	gttgggttta	atgggcattc	cttcttctgg	tcaaaaaac	60
276 gtcactggg	cacagaactt	atgttgttct	ctatggagaa	ctaaaagtat	gaggcgtagg	120
278 acactatttt	aattattttt	aatttattaa	tattaaata	tgtgaagctg	agtaattt	180
280 tctaagtcat	atttatattt	ttaagaagta	ccacttggaa	cattttatgt	attagtttg	240
282 aaataataat	ggaaagtggc	tatgcagtt	gaatatcctt	tgtttcagag	ccagatcatt	300
284 tcttggaaag	tgttagctta	cctcaaataa	atggctaact	tatacatatt	tttaaaagaaa	360
286 tatttatatt	gtatttatat	aatgtataaa	tggttttat	accaataat	ggcattttaa	420
288 aaaaattc						427

291 <210> SEQ ID NO: 16  
 292 <211> LENGTH: 15      *OK*  
 293 <212> TYPE RNA  
 294 <213> ORGANISM: Artificial Sequence  
 296 <220> FEATURE:

*Do you mean "u"?*      15

*see p. 6*

RAW SEQUENCE LISTING DATE: 07/28/2005  
 PATENT APPLICATION: US/10/543,033 TIME: 17:04:05

Input Set : D:\seq listing 10589-012-999 (as filed).txt  
 Output Set: N:\CRF4\07282005\J543033.raw

297 <223> OTHER INFORMATION: Description of Artificial Sequence: Group III AU-Rich element (ARE)

298 cluster of 3'untranslated region  
 301 <220> FEATURE:  
 302 <221> NAME/KEY: misc\_feature      *not allowed in an RNA sequence*  
 303 <222> LOCATION: (1)..(15)  
 304 <223> OTHER INFORMATION: n = a, t, g or c  
 306 <400> SEQUENCE: 16  
 W--> 307 nauuuauuu uuuuu      *same error*      15  
 310 <210> SEQ ID NO: 17  
 311 <211> LENGTH: 701      *in sequence 20*  
 312 <212> TYPE: DNA  
 313 <213> ORGANISM: Homo sapiens  
 315 <400> SEQUENCE: 17  
 316 aagagctcca gagagaagtc gaggaagaga gagacgggt cagagagac gcgcggcgt      60  
 318 gcgagcagcg aaagcgacag gggcaaagtg agtgcacgtc ttttgggggt gaccgcccga      120  
 320 gcgcggcgtg agccctcccc cttggatcc cgcaactgac cagtcgcgt gacggacaga      180  
 322 cagacagaca ccccccag ccccaattac cacccctcc cccggccggc gcgacagt      240  
 324 gacgcggcgg cgagccggg gcaggggccg gagccccc ccggaggccg gttggagggg      300  
 326 gtcggagctc gggcgctgc actgaaactt ttcttccaaac ttctgggtt ttctcgcttc      360  
 328 ggaggagccg tggccgcgc gggggaaagcc gagccgagcg gagccgcgag aagtgcgtac      420  
 330 tcggggccggg aggagccgca gccggaggag ggggaggagg aagaagagaa ggaagaggag      480  
 332 agggggccgc agtggcgact cggcgctcg aagccggct catggacggg tgaggccggc      540  
 334 gtgtgcgcag acagtgcgtcc agcgccgcgc ctcccccagcc ctggccggc ctcggccgg      600  
 336 gaggaagagt agctgcgcga ggcgcggagg agacggggcc gccccacagc cgcgcggga      660  
 338 gagggacgcg agccgcgcgc cccggtcggg cctccgaaac c      701  
 341 <210> SEQ ID NO: 18  
 342 <211> LENGTH: 1892  
 343 <212> TYPE: DNA  
 344 <213> ORGANISM: Homo sapiens  
 346 <400> SEQUENCE: 18  
 347 tgagccggc aggaggaagg agcctccctc agggtttcgg gaaccagatc tctctccagg      60  
 349 aaagactgat acagaacgat cgatacagaa accacgctgc cgccaccaca ccatcaccat      120  
 351 cgacagaaca gtcctaatac cagaaacctg aaatgaagga agaggagact ctgcgcagag      180  
 353 cactttgggt ccggagggcg agactccgc ggaagcattc cccggccgggt gaccagcac      240  
 355 gttcccttctt ggaattggat tcgcatttt attttcttg ctgtaataatc accgagcccg      300  
 357 gaagattaga gagtttatt tctggatcc ctgttagacac acccaccac atacatacat      360  
 359 ttatatatat atatattata tatataaaaa aataaatatc tctatttat atatataaaa      420  
 361 tatatatattt cttttttaa attaacatgt ctaatgttat tggtgtcttc actggatgt      480  
 363 ttgactgct gtggacttga gttggaggg gaatgttccc actcagatcc tgacagggaa      540  
 365 gaggaggaga tgagagactc tggcatgatc tttttttgt cccacttggt gggccaggg      600  
 367 tcctctcccc tggccaaagaa tgtgcaaggc cagggcatgg gggcaaatat gaccagtt      660  
 369 tggaaacacc gacaaaccca gccctggcgc tgagcccttc taccggatgt cagacggaca      720  
 371 gaaagacaaa tcacaggttc cgggatgagg acacggctc tgaccaggag tttggggagc      780  
 373 ttcaggacat tgcttgctt tggggattcc ctccacatgc tgacgcgcga tctcgccccc      840  
 375 aggggcactg cctgaaagat tcaggagct gggccgcctt cgcttactct cacctgcttc      900  
 377 ttagttgccc aggagccac tggcagatgt cccggcgaag agaagagaca cattgttgg      960  
 379 agaagcagcc catgacagcg ccccttcctg ggactcgcgc tcatcctt cctgctcccc      1020  
 381 ttctgggtt gcagctaaa aggacctatg tcctcacacc attgaaacca ctgttctgt      1080  
 383 ccccccagga aacctgggtt tgggtgttg agtgggtgac cttccat cccctggc      1140

see p. 7

10/543,033 7

<210> 40  
<211> 751  
<212> DNA  
<213> Homo sapiens

<220>  
<221> misc\_feature  
<222> (535). (739) " + " at location 739  
<223> n = a, t, g or c

<400> 40  
taagcaggcc tccaacgccc ctgtggccaa ctgcaaaaaa agcctccaag ggtttcgact 60  
ggtccagctc tgacatccct tcctggaaac agcatgaata aaacactcat cccatgggc 120  
caaattaata tgattctgct ccccccttct ccttttagac atggttgtgg gtctggaggg 180  
agacgtgggt ccaaggtcct catcccatcc tccctctgcc aggcaactatg tgtctggggc 240  
ttcgatcctt gggtgtcaggc agggctggga cacgcggctt ccctcccagt ccctgccttg 300  
gcaccgtcac agatgccaag caggcagcac ttagggatct cccagctggg ttagggcagg 360  
gcctggaaat gtgcattttg cagaaacttt tgagggtcgt tgcaagactg tgttagcaggc 420  
ctaccaggtc ccttcatct tgagagggac atggccctt gtttctgca gttccacgc 480  
ctctgcactc cctgccccctg gcaagtgcctc ccatcgcccc cggtgcccac catgnagctc 540  
cccgcacctg actcccccca catccaaggg cagccctgga accagtgggc tagttccttg 600  
aaggaagccc cactcattcc tattaatccc tcagaattcc cggggggagc cttccctct 660  
gaaccttgggt aaaaaatggg gaacgagaaa aaccccccgt tggagctgtg cgtttccagc 720  
ccctacttga gag~~not~~ttttt tttgggggccc g 751

see p 8

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<210> 88  
<211> 22  
<212> DNA  
<213> Artificial

Sequence  
sequence

<220>

<223> Description of Artificial Sequence: PCR primer (Sense/BglII)

RAW SEQUENCE LISTING ERROR SUMMARY                   DATE: 07/28/2005  
PATENT APPLICATION: US/10/543,033                   TIME: 17:04:06

Input Set : D:\seq listing 10589-012-999 (as filed).txt  
Output Set: N:\CRF4\07282005\J543033.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:1; N Pos. 3,7,8,11  
Seq#:2; N Pos. 3,4,7,8,11,12  
Seq#:3; N Pos. 3,4,7,8,11,12  
Seq#:13; N Pos. 1,15  
Seq#:16; N Pos. 1,15  
Seq#:20; N Pos..1,15  
Seq#:33; N Pos. 409,444  
Seq#:40; N Pos. 535,734

**VERIFICATION SUMMARY**  
PATENT APPLICATION: US/10/543,033

DATE: 07/28/2005  
TIME: 17:04:06

Input Set : D:\seq listing 10589-012-999 (as filed).txt  
Output Set: N:\CRF4\07282005\J543033.raw

L:11 M:270 C: Current Application Number differs, Replaced Application Number  
L:12 M:271 C: Current Filing Date differs, Replaced Current Filing Date  
L:45 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:0  
L:70 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:0  
L:95 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:0  
L:254 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13 after pos.:0  
L:307 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:16 after pos.:0  
L:445 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:20 after pos.:0  
L:1078 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:33 after pos.:360  
M:341 Repeated in SeqNo=33  
L:1289 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:40 after pos.:480  
M:341 Repeated in SeqNo=40  
L:2695 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:88